SEQUENCE LISTING

<110> Trimeris, Inc.

<120> HIV-Derived HR1 Peptides Modified to Form Stable Trimers, and Their Use In Therapy to Inhibit Transmission of Human Immunodeficiency Virus

<130> TRM-001

<150> US 60/414,514 <151> 2002-09-27

<160> 82

<170> PatentIn version 3.2

<210> 1 <211> 59 <212> PRT

<213> Human immunodeficiency virus type 1

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Leu Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val 35 40

Glu Arg Tyr Leu Lys Asp Gln Leu Leu Gly Ile

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Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu

Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu

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 Gln Leu Gln Ala Arg Ile
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Gln His Leu Leu
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Gly Ile Val Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln
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Gln His Leu Leu Gln Leu
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Gly Ile Val Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln
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Gln His Leu Ceu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln Ala
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Arg Ile
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Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln Leu Leu Ser Gly
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Ile Val Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln
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His Leu Leu Gln
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Val Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His
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Leu Leu Gln Leu
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Ser Met Thr Leu Thr Val Gln Ala Arg Gln Leu Leu Ser Gly Ile Val
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Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu
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Leu Gln Leu Thr
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Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu
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Gln Leu Thr
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Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu
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Gln Leu Thr Val
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Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln 20 25 30

Leu Thr Val

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Thr Leu Thr Val Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln 1 5 10 15

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Leu Thr Val Trp 35

<210> 17

<211> 37

<212> PRT

<213> Artificial

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<223> synthesized

<400> 17

Thr Leu Thr Val Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln 1 5 10 15

Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln 20 25 30

Leu Thr Val Trp Gly 35

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Leu Thr Val Trp Gly Ile
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Thr Leu Thr Val Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln
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Leu Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg
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Leu Thr Val Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Gln
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Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu
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Thr Val Trp Gly
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Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Asn Asn Leu
                                     10
                 5
 Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu Thr Val Trp
 Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu
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  Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu Thr Val Trp
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  Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr
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Leu Arg Ala Ile Glu Ala Gln Gln His Leu Gln Leu Thr Val Trp
                                25
Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu
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Lys
      24
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Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Gln Asn Asn Leu
Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu Thr Val Trp
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                                25
Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu
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Lys Asp Gln
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Ser Gly Ile Val Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala 1 5 10 15

Gln Gln His Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln 20 25 30

Ala Arg Ile Leu 35

<210> 26

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<212> PRT

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<220>

<223> synthesized

<400> 26

Ser Gly Ile Val Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala 1 5 10 15

Gln Gln His Leu Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln 20 25 30

Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu Lys Asp Gln 35 40 45

<210> 27

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<212> PRT

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<400> 27

Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu 20 25 30

Ala Val Glu Arg Tyr Leu Lys Asp Gln 35 40

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<210> 28
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Arg Ala Ile Glu Ala Gln Gln His Leu Cln Leu Thr Val Trp Gly
                                    10
                5
Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu Lys
                                                    30
                                25
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Asp Gln
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Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu
                5
Leu Gln Leu Thr Ala Trp Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu
                                25
            20
Ala Val Glu Arg Tyr Leu Lys Asp Gln
        35
<210> 30
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Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu
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Leu Gln Leu Thr Val Ala Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu 20 25 30

Ala Val Glu Arg Tyr Leu Lys Asp Gln 35 40

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Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Asn Asn Leu 1 5 10 15

Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu Thr Val Phe 20 25 30

Gly Ile Arg Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu $35 \hspace{1cm} 40 \hspace{1cm} 45$

Lys

<210> 32

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<220>

<223> synthesized

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Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Gln Asn Asn Leu 1 5 10 15

Leu Arg Ala Ile Glu Ala Gln Gln His Ala Leu Gln Ala Thr Val Trp 20 25 30

Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu 35 40 45

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<210> 35 <211> 49 <212> PRT <213> Artificial

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Gln Ala Arg Gln Leu Val Ser Gly Leu Val Gln Gln Asn Asn Ile
Leu Arg Ala Leu Glu Ala Thr Gln His Ala Val Gln Ala Leu Val Trp
Gly Val Lys Gln Leu Gln Ala Arg Val Leu Ala Leu Glu Arg Tyr Ile
                           40
Lys
<210> 36
<211> 49
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Gln Ile Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Asn Asn Leu
                                                       15
                5
Leu Arg Ala Ile Glu Ala Ile Gln His Ala Leu Gln Ala Ile Val Trp
Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu
Lys
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<400> 37

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Gln Ala Arg Gln Leu Val Ser Gly Leu Val Gln Gln Asn Asn Ile
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Gly Val Arg Gln Leu Gln Ala Arg Val Leu Ala Leu Glu Arg Tyr Ile
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  Leu Arg Ala Ile Glu Ala Thr Gln His Ala Val Gln Ala Leu Val Trp
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   Gly Val Lys Gln Leu Gln Ala Arg Val Leu Ala Leu Glu Arg Tyr Ile
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   Lys Asp Gln
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   <210> 39
    <211> 51
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    Gln Ala Arg Gln Leu Val Ser Gly Leu Val Gln Gln Asn Asn Ile
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Leu Arg Ala Leu Glu Ala Gln Gln His Ala Leu Gln Ala Thr Val Trp 20 25 30

Gly Ile Lys Gln Leu Gln Ala Arg Val Leu Ala Leu Glu Arg Tyr Ile 35 40 45

Lys Asp Gln 50

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Gln Ala Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Gln Asn Asn Leu 1 5 10 15

Leu Arg Ala Ile Glu Ala Gln Gln His Ala Leu Gln Ala Thr Val Trp 20 25 30

Gly Val Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu 35 40 45

Lys Asp Gln 50

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Gln Ala Arg Gln Leu Val Ser Gly Leu Val Gln Gln Gln Asn Ile

5 10 15

Leu Arg Ala Leu Glu Ala Thr Gln His Leu Val Gln Leu Leu Val Trp 20 25 30

Gly Val Lys Gln Leu Gln Ala Arg Val Leu Ala Leu Glu Arg Tyr Ile

35 40 45

Lys

<210> 42

<211> 49

<212> PRT

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Gln Ile Arg Gln Leu Leu Ser Gly Ile Val Gln Gln Asn Asn Leu 1 5 10 15

Leu Arg Ala Ile Glu Ala Ile Gln His Leu Leu Gln Leu Ile Val Trp 20 25 30

Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$

Lys

<210> 43

<211> 41

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Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Leu Gln Leu Thr Val Phe Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu 20 25 30

Ala Val Glu Arg Tyr Leu Lys Asp Gln 35 40

<210> 44

<211> 41

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Leu Gln Leu Thr Val Trp Gly Ile Ala Gln Leu Gln Ala Arg Ile Leu
Ala Val Glu Arg Tyr Leu Lys Asp Gln
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Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Ala Ala Arg Ile Leu
                               25
Ala Val Glu Arg Tyr Leu Lys Asp Gln
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Gln Gln Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Ala
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Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu

5

15

20 25 30

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atcctg	gctg ttgaacgtta	cctgaaa				147
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gttctgg	yctc tggaacgtta	catcaaa				147
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cagatco	egte agetgetgte	tggtatcgtt	cagcagcaga	acaacctgct	gcgtgctatc	60
gaagcta	atcc agcacgctct	gcaggctatc	gtttggggta	tcaaacagct	gcaggctcgt	120
atcctgg	gctg ttgaacgtta	cctgaaa				147
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cag							123
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cag							123
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gaggccc	cagc	agcacgccct	gcaggccacc	gtgtggggca	tcaagcagct	gcaggcccgc	120
atcctgg	geeg	tggagcgcta	cctgaag				147
	53 147 DNA Arti	ficial					
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atcctg	gccg tggagcgcta	cctgaag				147
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cag						123
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gtggcc	ggca tcaagcagct	gcaggcccgc	atcctggccg	tggagcgcta	cctgaaggac	120
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<210><211><212><212><213>	57 8 PRT Artificial					

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Trp Xaa Xaa Trp Xaa Xaa Xaa Ile
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Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu Arg Tyr Leu 40

Lys Asp Gln 50

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